# Name:

# **Enrolment No:**



# **UPES**

# **End Semester Examination, December 2024**

Course: Systems Biology
Program: B.Tech Biotechnology
Course Code: HSBT3005
Semester : V
Duration : 3 Hours
Max. Marks: 100

**Instructions: Attempt all questions** 

S. No.	Section A	Marks	COs
	Short answer questions/ MCQ/T&F		
	(20Qx1.5M= 30 Marks)		
Q 1	Which omics technology is most suitable for studying post-translational modifications?	1.5	CO1
	a) Metabolomics		
	b) Proteomics		
	c) Genomics		
	d) Lipidomics		
Q 2	The rate of an enzyme-catalyzed reaction is directly proportional to:	1.5	CO1
	a) Substrate concentration		
	b) Enzyme concentration		
	c) Both enzyme and substrate concentration		
	d) Product concentration		
Q 3	Which metabolic pathway synthesizes glucose from non-	1.5	CO1
	carbohydrate sources?		
	a) Glycogenesis		
	b) Glycolysis		
	c) Gluconeogenesis		
	d) Pentose phosphate pathway		
Q 4	Gene expression can be regulated post-transcriptionally by:	1.5	CO1
	a) Promoter methylation		
	b) RNA interference		
	c) DNA methylation		
	d) Histone acetylation		
Q 5	Describe the role of feedback inhibition in metabolic pathways.	1.5	CO1
Q 6	Pathway enrichment analysis is used to:	1.5	CO2
	a) Annotate non-coding regions		
	b) Identify over-represented pathways in datasets		
	c) Sequence whole genomes		
	d) Compare protein domains		
Q 7	Which tool is widely used for genome annotation?	1.5	CO2

	a) DI ACT		
	a) BLAST		
	b) RAST		
	c) CRISPR/Cas9		
	d) ClustalW	1.5	002
Q 8	Why do extremophiles have unique metabolic pathways?	1.5	CO2
	a) To enhance photosynthesis		
	b) To adapt to extreme environments		
	c) To reduce energy requirements		
	d) To perform horizontal gene transfer		
Q 9	Metabolic pathways in halophiles often involve:	1.5	CO2
	a) Reduced osmotic stress		
	b) Increased glycolysis rates		
	c) Specialized ion transport mechanisms		
	d) Anaerobic respiration		
Q 10	The study of organism-specific metabolic pathways can help in:	1.5	CO2
	a) Constructing phylogenetic trees		
	b) Identifying novel enzymes and biocatalysts		
	c) Developing RNA vaccines		
	d) Determining ribosome structures		
Q 11	Metabolic control analysis (MCA) focuses on identifying:	1.5	CO3
	a) Pathways involved in cell division		
	b) Key enzymes regulating metabolic flux		
	c) The structure of DNA		
	d) Gene expression patterns		
Q 12	In Drosophila melanogaster, the regulatory network is crucial for:	1.5	CO3
	a) Protein folding		
	b) Developmental processes		
	c) Metabolite transport		
	d) Gene expression regulation		
Q 13	The MAP kinase cascade is an example of a:	1.5	CO3
	a) Metabolic pathway		
	b) Signaling pathway		
	c) Genetic network		
	d) Protein interaction network		
Q 14	Which of the following best describes a mechanical network in cell	1.5	CO3
	biology?		
	a) A network of cell signaling pathways		
	b) A network governing the cytoskeleton dynamics		
	c) A network of enzyme interactions		
	d) A network involving metabolic reactions		
Q 15	The key assumption in flux balance analysis is:	1.5	CO3
415	a) The system reaches equilibrium at steady-state	1.5	
	b) All reactions are reversible		
	c) Enzyme concentrations are uniform		
	c) Enzyme concentrations are uniform		

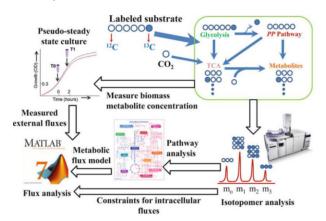
	d) Gene expression is linear					
0.16	-	1.5	CO4			
Q 16	What happens in the lac operon when lactose is present in the	1.5	CO4			
	environment?					
	a) The repressor binds to the operator					
	b) The repressor is inactivated and transcription proceeds					
	c) The RNA polymerase is inhibited					
	d) The gene for the repressor is activated					
Q 17	In systems biology, which approach is used to model large-scale	1.5	CO4			
	gene regulatory networks?					
	a) Stochastic modeling					
	b) Linear regression					
	c) Genome-scale metabolic modeling					
	d) Principal component analysis					
Q 18	Which of the following is an open-source tool for modeling	1.5	CO4			
	biological systems?					
	a) MATLAB					
	b) Ecell					
	c) Excel					
	d) AutoCAD					
Q 19	BioNets is used for:	1.5	CO4			
	a) Protein structure analysis					
	b) Modeling gene regulatory networks					
	c) Gene expression profiling					
	d) Metabolic pathway analysis					
Q 20	What is the purpose of the E. coli chemotactic pathway?	1.5	CO4			
220	a) To promote DNA repair	1.0				
	b) To enable movement toward attractants and away from					
	repellents					
	c) To regulate protein synthesis					
	d) To control bacterial growth rate					
	-					
	Section B: Short-Answer Questions (4Qx5M=20 Marks)					
	(4QX5W1-20 Walks)					
Q 1	Define systems biology? Discuss its importance in predicting	5	CO1			
~ 1	phenotypic behavior in an organism using omics data.					
Q 2	Discuss the role genome annotation plays in predicting metabolic	5	CO2			
42	pathways? Explain the importance of functional gene annotation.		002			
Q 3	List various factors affecting distribution of fluxes across metabolic	5	CO3			
\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \		, ,				
0.4	pathways  Describe the primary function of the DAVID software in functional	5	CO4			
Q 4		)	004			
	genomics, and how does it assist in biological research?					
Section C: Case study						
	South St. Sube State					

## (2Qx15M=30 Marks)

The goal of metabolic engineering is to design and build engineered biological systems that can produce chemicals, materials, food, and drugs at high yield using the appropriate microorganisms. However, the lack of fundamental understanding of cellular responses during industrial bioprocesses often prevents metabolic engineers from achieving satisfactory goals in biochemical production. In the past decade, 13C-MFA has been widely used to provide insightful information on metabolism of various microorganisms, thus helping metabolic engineers to successfully improve biochemical production.

Q 1

Stable isotope labeling is a powerful technique with promising applications. Depending upon this powerful technology, labeled tracers can sensitively and accurately track changes according to the location and quantity of peptides, amino acids, or carbohydrates containing isotope-labeled in vivo or in vitro. It enables direct analysis of nutrient distribution, metabolism, conversion into metabolites, and the fate of the resulting metabolites. In contrast to radioactive labeling, there are no dangers or safety concerns, making this technique particularly well suited for metabolism studies in humans. As a result, isotope labeling technology has received progressively more recognition in the fields of medicine and biochemistry.



Look at the diagram carefully and answer the following:

- A) Which approach is depicted in the diagram to estimating metabolic fluxes for biochemical production. Explain this approach and its significance in metabolic engineering.
- B) With the help of the given diagram, list five key procedural steps in estimating the metabolic fluxes
- C) Define isotopomers? Why is isotopic labelling important?

15 marks (3 marks each)

	D) Explain isotopomer analysis and two primary analytical		
	methods used for measuring the labelled metabolites.		
	E) List any two promising applications of stable isotope		
	labeling.		
Q 2	A biotech startup is developing a novel gene therapy for the	15 marks	CO4
	treatment of rare genetic disorders. The startup uses gene editing	(5 marks	
	technologies, like CRISPR-Cas9, and combines them with systems	each)	
	biology to model the effects of gene modification on cellular		
	pathways. They aim to personalize gene therapies based on the		
	patient's genetic profile, ensuring more effective and safer		
	treatments.		
	Based on case study, answer the following:		
	A) Discuss the significance of gene editing technologies like		
	CRISPR-Cas9 in biotechnology? How does systems biology		
	contribute to the development of personalized gene		
	therapies?		
	B) What challenges are faced when developing gene therapies		
	for rare genetic disorders? Discuss the role of patient genetic		
	profiling in personalized gene therapy.		
	C) Which ethical considerations must be addressed when using		
	gene editing technologies in gene therapy?		
	g		
	Section D: Long-Answer Questions		
	(2Qx10M=20 Marks)		
Q 1	A) Explain the significance of promoter regions in genome	5+5	CO3
	regulation. How do mutations in promoter regions affect gene	marks	
	expression?		
	B) Explain the role of mRNA stability in post-transcriptional		
	regulation. What factors influence mRNA degradation?		
Q 2	Explain the importance of mathematical models in representing cell	10 marks	CO4
	biological systems. How can metabolic network analysis be used to		
	identify drug targets in pathogenic bacteria?		